```
<110> Lal, Preeti
Faris, Mary
Chen, Huei-Mei
Ison, Craig H.
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<120> STEAP-RELATED PROTEIN

<130> PC-0037 US

<140> To Be Assigned

<141> Herewith

<160> 11

<170> PERL Program

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<212> PRT

<213> Homo sapiens

<220>

<221> misc\_feature

<223> Incyte ID No: 7492448CD1

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                 35
                                     40
Thr Ile Arg Leu Ile Arg Cys Gly Tyr His Val Val Ile Gly Ser
                 50
                                     55
Arg Asn Pro Lys Phe Ala Ser Glu Phe Phe Pro His Val Val Asp
                 65
                                     70
Val Thr His His Glu Asp Ala Leu Thr Lys Thr Asn Ile Ile Phe
                 80
                                     85
Val Ala Ile His Arg Glu His Tyr Thr Ser Leu Trp Asp Leu Arg
                 95
                                    100
His Leu Leu Val Gly Lys Ile Leu Ile Asp Val Ser Asn Asn Met
                110
                                    115
Arg Ile Asn Gln Tyr Pro Glu Ser Asn Ala Glu Tyr Leu Ala Ser
                125
                                     130
                                                         135
Leu Phe Pro Asp Ser Leu Ile Val Lys Gly Phe Asn Val Val Ser
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                                    145
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Ala Trp Ala Leu Gln Leu Gly Pro Lys Asp Ala Ser Arg Gln Val
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                                    160
Tyr Ile Cys Ser Asn Asn Ile Gln Ala Arg Gln Gln Val Ile Glu
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                                    175
Leu Ala Arg Gln Leu Asn Phe Ile Pro Ile Asp Leu Gly Ser Leu
                185
                                    190
                                                         195
Ser Ser Ala Arg Glu Ile Glu Asn Leu Pro Leu Arg Leu Phe Thr
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Leu Trp Arg Gly Pro Val Val Val Ala Ile Ser Leu Ala Thr Phe
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                                    220
                                                         225
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Met Glu Ser Ile Ser Met Met Gly Ser Pro Lys Ser Leu Ser Glu

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Phe Phe Leu Tyr Ser Phe Val Arg Asp Val Ile His Pro Tyr Ala
                230
                                     235
                                                          240
Arg Asn Gln Gln Ser Asp Phe Tyr Lys Ile Pro Ile Glu Ile Val
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                                                          255
Asn Lys Thr Leu Pro Ile Val Ala Ile Thr Leu Leu Ser Leu Val
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                                     265
                                                          270
Tyr Leu Ala Gly Leu Leu Ala Ala Ala Tyr Gln Leu Tyr Tyr Gly
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                                     280
                                                          285
Thr Lys Tyr Arg Arg Phe Pro Pro Trp Leu Glu Thr Trp Leu Gln
                290
                                     295
                                                          300
Cys Arg Lys Gln Leu Gly Leu Leu Ser Phe Phe Phe Ala Met Val
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                                     310
                                                          315
His Val Ala Tyr Ser Leu Cys Leu Pro Met Arg Arg Ser Glu Arg
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                                     325
                                                          330
Tyr Leu Phe Leu Asn Met Ala Tyr Gln Gln Val His Ala Asn Ile
                335
                                     340
                                                          345
Glu Asn Ser Trp Asn Glu Glu Glu Val Trp Arg Ile Glu Met Tyr
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                                                          360
                                     355
Ile Ser Phe Gly Ile Met Ser Leu Gly Leu Leu Ser Leu Leu Ala
                365
                                     370
                                                          375
Val Thr Ser Ile Pro Ser Val Ser Asn Ala Leu Asn Trp Arg Glu
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                                     385
                                                          390
Phe Ser Phe Ile Gln Ser Thr Leu Gly Tyr Val Ala Leu Leu Ile
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                                     400
                                                          405
Ser Thr Phe His Val Leu Ile Tyr Gly Trp Lys Arg Ala Phe Glu
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Glu Glu Tyr Tyr Arg Phe Tyr Thr Pro Pro Asn Phe Val Leu Ala
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Leu Val Leu Pro Ser Ile Val Ile Leu Gly Lys Ile Ile Leu Phe
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                                     445
Leu Pro Cys Ile Ser Arg Lys Leu Lys Arg Ile Lys Lys Gly Trp
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                                     460
                                                          465
Glu Lys Ser Gln Phe Leu Glu Glu Gly Ile Gly Gly Thr Ile Pro
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<223> Incyte ID No: 7492448CB1

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<sup>&</sup>lt;211> 1891

<sup>&</sup>lt;212> DNA

<sup>&</sup>lt;213> Homo sapiens

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tattcccaga ttctttgatt gtcaaaggat ttaatgttgt ctcagcttgg gcacttcagt 780
taggacctaa ggatgccagc cggcaggttt atatatgcag caacaatatt caagcgcgac 840
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gaggaacaat teeteatgte teeeeggaga gggteacagt aatgtgatga taaatggtgt 1800
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                                                                  1891
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<213> Homo sapiens

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PC-0037 US
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taaaatcctg attgatgtga gcaataacat gaggataaac cagtacccag aatccaatgc 300
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PC-0037 US
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tgcttccgat gaccacgtga tagccgcacc tgataagccg aatggtcaga gacttggcaa 180
aatccccact tcctatcacc cccacggtga ccttccttgc gtctttgata ccgtttatgc 240
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cagaggagct ggaggaggc tccttcggcg ctgctggact ctggaactgc ctacgtgtag 420
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Val Ser Asp Ser Leu Thr Trp Arg Glu Phe His Tyr Ile Gln Ser

Lys Leu Gly Ile Val Ser Leu Leu Gly Thr Ile His Ala Leu

235

250

255

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Ile	Phe	Ala	$\operatorname{Trp}$		Lys	Trp	Ile	Asp	Ile	Lys	Gln	Phe	Val	_
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Tyr	Thr	Pro	Pro	Thr	Phe	Met	Ile	Ala	Val	Phe	Leu	Pro	Ile	Val
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Val	Leu	Ile	Phe	Lys	Ser	Ile	Leu	Phe	Leu	Pro	Cys	Leu	Arg	Lys
				305					310					315
Lys	Ile	Leu	Lys	Ile	Arg	His	Gly	Trp	Glu	Asp	Val	Thr	Lys	Ile
				320					325				_	330
Asn	Lys	Thr	Glu	Ile	Cys	Ser	Gln	Leu						
	_			335										